



IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

PATENT

By the Application of:

REBECCA E. CAHOON ET AL.  
APPLN. NO.: 09/720,529  
FILED: DECEMBER 20, 2000  
FOR: CHROMATIN ASSOCIATED  
PROTEINS

CASE NO.: BB-1118-A  
GROUP ART UNIT: 1652  
EXAMINER: RICHARD G. HUTSON  
CONFIRMATION NO.: 2357

1652  
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M.G.J  
2/24/03

**PRELIMINARY AMENDMENT AND  
RESPONSE TO RESTRICTION REQUIREMENT**

**RECEIVED**

Assistant Commissioner for Patents  
Washington, DC 20231

FEB 19 2003  
TECH CENTER 1600/2900

Sir:

This is a Preliminary Amendment and Response to the Restriction Requirement set forth in the Office Action mailed November 5, 2002. A Petition for Extension of Time for two (2) months up to and including February 5, 2002, is filed simultaneously herewith. Please enter the following:

**IN THE SPECIFICATION**

Please amend the specification as follows; a marked-up version showing changes made is attached hereto:

**Paragraph beginning at page 4, line 36, and continuing through page 5, line 20:**

B' A "substantial portion" of an amino acid or nucleotide sequence comprises an amino acid or a nucleotide sequence that is sufficient to afford putative identification of the protein or gene that the amino acid or nucleotide sequence comprises. Amino acid and nucleotide sequences can be evaluated either manually by one skilled in the art, or by using computer-based sequence comparison and identification tools that employ algorithms such as BLAST (Basic Local Alignment Search Tool; Altschul et al. (1993) *J. Mol. Biol.* 215:403-410). In general, a sequence of ten or more contiguous amino acids or thirty or more contiguous nucleotides is necessary in order to putatively identify a polypeptide or nucleic acid sequence as homologous to a known protein or gene. Moreover, with respect to nucleotide sequences, gene-specific oligonucleotide probes comprising 30 or more contiguous nucleotides may be used in sequence-dependent methods of gene identification (e.g., Southern hybridization) and isolation (e.g., *in situ* hybridization of bacterial colonies or bacteriophage plaques). In addition, short oligonucleotides of 12 or more nucleotides